

SEQUENCE LISTING

<110> WATANABE, Eijiro OEDA, Kenji <120> Raffinose Synthase Genes and Their Use																	
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									gac Asp								163
									tcc Ser 30								211
	_			-					ttc Phe				-				259
									gct Ala								307
				_					aac Asn				_	_	_		355
									tcc Ser								403

451

gta gtt cca ctc ggc aaa cta aaa gga atc aaa ttc atg agc ata ttc Val Val Pro Leu Gly Lys Leu Lys Gly Ile Lys Phe Met Ser Ile Phe $\,$

105	110	115
105	110	113

			_							gtc Val					_	499
										ctg Leu						547
										atc Ile 160						595
_									_	cac His			_		_	643
_	_					_				agc Ser			_	_		691
										ata Ile						739
	_				_					aag Lys			-	_		787
	-		-			_				tgg Trp 240	_	_		_	-	835
		_	_	-					_	tgg Trp	_		-	_		883
		-			_					gtc Val			-	_		931
				-		-	_	_	-	gaa Glu	-	-			-	979
										tgc Cys						1027
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										ttg Leu						1123
										ctt Leu						1171

		agg Arg 360														1219
		gtg Val														1267
_	_	att Ile	_							-	_		-			1315
		atg Met														1363
-		gtt Val		-	-	_			_							1411
_		ggt Gly 440		_	_			-	_	-				_		1459
		tca Ser	-	_								_		-	-	1507
_		cat His	-		_							-	-			1555
		cgc Arg														1603
		ggt Gly							_		_	_		_	_	1651
tac Tyr	aac Asn	agt Ser 520	tta Leu	tgg Trp	atg Met	gga Gly	aat Asn 525	ttc Phe	att Ile	cag Gln	cca Pro	gat Asp 530	tgg Trp	gac Asp	atg Met	1699
		tcc Ser														1747
		ggc Gly					-									1795
	_	ttg Leu					-	_		_				_	_	1843
_		cat His		-				-	-	_	_		-	-		1891

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		_		ggg tgg tgt Gly Trp Cys	
	-	-	_	cgc gcg gtg Arg Ala Val	
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				gat aat gca Asp Asn Ala 740	
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				tgg cct agt Trp Pro Ser	
		cag ttt tta Gln Phe Leu		tag gaatcctat	.g 2517
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Leu Lys Glu Asn Thr Phe Arg Thr Ser Leu Gln Pro Gly Leu Asn Asp His 175

Ile Gly Met Ser Val Glu Ser Gly Ser Thr His Val Thr Gly Ser Ser 180

Phe Lys Ala Cys Leu Tyr Ile His Leu Ser Asn Asp Pro Tyr Ser Ile 200

Leu Lys Glu Ala Val Lys Val Ile Gln Thr Gln Leu Gly Thr Phe Lys 210

Thr Leu Glu Glu Lys Thr Ala Pro Ser Ile Ile Asp Lys Phe Gly Trp 225 230 235 240

Cys Thr Trp Asp Ala Phe Tyr Leu Lys Val His Pro Lys Gly Val Trp 245 250 255

Glu Gly Val Lys Ser Leu Thr Asp Gly Gly Cys Pro Pro Gly Phe Val 260 265 270

Ile Ile Asp Asp Gly Trp Gln Ser Ile Cys His Asp Asp Asp Glu 275 280 285

Asp Asp Ser Gly Met Asn Arg Thr Ser Ala Gly Glu Gln Met Pro Cys

290	295	300
200	233	500

305	Leu	Val	Lys	Tyr	Glu 310	Glu	Asn	Ser	Lys	Phe 315	Arg	Glu	Tyr	Glu	Asn 320
Pro	Glu	Asn	Gly	Gly 325	Lys	Lys	Gly	Leu	Gly 330	Gly	Phe	Val	Arg	Asp 335	Leu
Lys	Glu	Glu	Phe 340	Gly	Ser	Val	Glu	Ser 345	Val	Tyr	Val	Trp	His 350	Ala	Leu
Cys	Gly	Tyr 355	Trp	Gly	Gly	Val	Arg 360	Pro	Gly	Val	His	Gly 365	Met	Pro	Lys
Ala	Arg 370	Val	Val	Val	Pro	Lys 375	Val	Ser	Gln	Gly	Leu 380	Lys	Met	Thr	Met
Glu 385	Asp	Leu	Ala	Val	Asp 390	Lys	Ile	Val	Glu	Asn 395	Gly	Val	Gly	Leu	Val 400
Pro	Pro	Asp	Phe	Ala 405	His	Glu	Met	Phe	Asp 410	Gly	Leu	His	Ser	His 415	Leu
Glu	Ser	Ala	Gly 420	Ile	Asp	Gly	Val	Lys 425	Val	Asp	Val	Ile	His 430	Leu	Leu
Glu	Leu	Leu 435	Ser	Glu	Glu	Tyr	Gly 440	Gly	Arg	Val	Glu	Leu 445	Ala	Arg	Ala
Туr	Tyr 450	Lys	Ala	Leu	Thr	Ser 455	Ser	Val	Lys	Lys	His 460	Phe	Lys	Gly	Asn
Gly	Val	Ile	Ala	Ser	Met	Glu	His	Cys	Asn	Asp	Phe	Phe	Leu	Leu	Gly
465					470					475					480
465	Glu	Ala	Ile	Ser 485		Gly	Arg	Val	Gly 490		Asp	Phe	Trp	Cys 495	
465 Thr	Glu Pro			485	Leu				490	Asp				495	Ser
465 Thr Asp		Ser	Gly 500	485 Asp	Leu Pro	Asn	Gly	Thr 505	490 Tyr	Asp Trp	Leu	Gln	Gly 510	495 Cys	Ser His
465 Thr Asp Met	Pro	Ser His 515	Gly 500 Cys	Asp Ala	Leu Pro Tyr	Asn Asn	Gly Ser 520	Thr 505 Leu	490 Tyr Trp	Asp Trp Met	Leu Gly	Gln Asn 525	Gly 510 Phe	495 Cys Ile	Ser His Gln
465 Thr Asp Met	Pro Val Asp	Ser His 515 Trp	Gly 500 Cys Asp	485 Asp Ala Met	Leu Pro Tyr Phe	Asn Asn Gln 535	Gly Ser 520 Ser	Thr 505 Leu Thr	490 Tyr Trp His	Asp Trp Met Pro	Leu Gly Cys 540	Gln Asn 525 Ala	Gly 510 Phe Glu	495 Cys Ile Phe	Ser His Gln His
465 Thr Asp Met Pro Ala 545	Pro Val Asp 530	Ser His 515 Trp Ser	Gly 500 Cys Asp	Asp Ala Met	Leu Pro Tyr Phe Ile 550	Asn Asn Gln 535 Ser	Gly Ser 520 Ser	Thr 505 Leu Thr	490 Tyr Trp His	Asp Trp Met Pro Ile 555	Leu Gly Cys 540	Gln Asn 525 Ala Val	Gly 510 Phe Glu Ser	495 Cys Ile Phe Asp	Ser His Gln His Cys 560
A65 Thr Asp Met Pro Ala 545 Val	Pro Val Asp 530 Ala	Ser His 515 Trp Ser Asn	Gly 500 Cys Asp Arg	Asp Ala Met Ala Asn 565	Leu Pro Tyr Phe Ile 550 Phe	Asn Asn Gln 535 Ser Lys	Gly Ser 520 Ser Gly Leu	Thr 505 Leu Thr Gly	490 Tyr Trp His Pro	Asp Trp Met Pro Ile 555 Ser	Leu Gly Cys 540 Tyr	Gln Asn 525 Ala Val	Gly 510 Phe Glu Ser Leu	495 Cys Ile Phe Asp	Ser His Gln His Cys 560 Asp
A65 Thr Asp Met Pro Ala 545 Val	Pro Val Asp 530 Ala Gly	Ser His 515 Trp Ser Asn	Gly 500 Cys Asp Arg His	Asp Ala Met Ala Asn 565 Arg	Leu Pro Tyr Phe Ile 550 Phe Cys	Asn Asn Gln 535 Ser Lys	Gly Ser 520 Ser Gly Leu	Thr 505 Leu Thr Gly Leu Tyr 585	490 Tyr Trp His Pro Lys 570 Ala	Asp Trp Met Pro Ile 555 Ser Leu	Leu Gly Cys 540 Tyr Leu Pro	Gln Asn 525 Ala Val Val	Gly 510 Phe Glu Ser Leu Arg 590	495 Cys Ile Phe Asp Pro 575 Asp	Ser His Gln His Cys 560 Asp Cys

Gly 625	Gly	Trp	Cys	Pro	Glu 630	Ala	Arg	Arg	Asn	Lys 635	Ser	Val	Ser	Glu	Phe 640	
Ser	Arg	Ala	Val	Thr 645	Cys	Tyr	Ala	Ser	Pro 650	Glu	Asp	Ile	Glu	Trp 655	Cys	
Asn	Gly	Lys	Thr 660	Pro	Met	Ser	Thr	Lys 665	Gly	Val	Asp	Phe	Phe 670	Ala	Val	
Tyr	Phe	Phe 675	Lys	Glu	Lys	Lys	Leu 680	Arg	Leu	Met	Lys	Cys 685	Ser	Asp	Arg	
Leu	Lys 690	Val	Ser	Leu	Glu	Pro 695	Phe	Ser	Phe	Glu	Leu 700	Met	Thr	Val	Ser	
Pro 705	Val	Lys	Val	Phe	Ser 710	Lys	Arg	Phe	Ile	Gln 715	Phe	Ala	Pro	Ile	Gly 720	
Leu	Val	Asn	Met	Leu 725	Asn	Ser	Gly	Gly	Ala 730	Ile	Gln	Ser	Leu	Glu 735	Phe	
Asp	Asp	Asn	Ala 740	Ser	Leu	Val	Lys	Ile 745	Gly	Val	Arg	Gly	Cys 750	Gly	Glu	
Met	Ser	Val 755	Phe	Ala	Ser	Glu	Lys 760	Pro	Val	Cys	Cys	Lys 765	Ile	Asp	Gly	
Val	Lys 770	Val	Lys	Phe	Leu	Tyr 775	Glu	Asp	Lys	Met	Ala 780	Arg	Val	Gln	Ile	
Leu 785	Trp	Pro	Ser	Ser	Ser 790	Thr	Leu	Ser	Leu	Val 795	Gln	Phe	Leu	Phe		
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			-	-	-	-		_	al Gl					ne G	gt ctt Ly Leu L5	109
-	aac Asn			_		_					-					157
	gcc Ala															205
_	acc Thr					_	_	_	_	_	_				-	253

50				55				60					
gac gac Asp Asp 65	-		_		 				-			_	301
agc cga Ser Arg	_			-	 _		_					-	349
agc ata Ser Ile	Phe A												397
aac gga Asn Gly	_	_				_	_				_		445
aac gac Asn Asp 130	Gln L		Arg			_			_				493
gcc tcg Ala Ser 145				-			_	_	_			-	541
gtt tgc Val Cys				_	_	-	-			_			589
agc tgc Ser Cys	Leu T	-		_		_	_		_	_		_	637

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					ctg Leu											1165
			-	_	atc Ile	_	_			-		_				1213
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					gtt Val											1309
					ggt Gly											1357
					tcg Ser											1405
					cat His											1453
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		_			ggc Gly	_						_		_		1549
	-	-			agc Ser	_		_					_	_	_	1597
	_	_		_	tcc Ser				_	_	-			_	_	1645
		-			ggt Gly			-		-	_	_	-	-		1693

								agc Ser								1741
							-	ctc Leu			_	-	_	-		1789
_	-		_		-		_	aca Thr 585	_							1837
								cta Leu			_					1885
								aag Lys								1933
								caa Gln								1981
	_			_				atg Met				-	-		_	2029
	_	_				_		atg Met 665	_	_				_	_	2077
_								gag Glu		_						2125
		_			_			caa Gln		_						2173
								att Ile								2221
								gtt Val								2269
								agt Ser 745								2317
								atg Met								2365
cct Pro	agt Ser 770	gct Ala	tca Ser	aaa Lys	ttg Leu	tca Ser 775	atg Met	gtt Val	gag Glu	ttt Phe	tta Leu 780	ttt Phe	tgat	ccct	:ga	2414

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2498

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(215) Grycine me

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Val Asn Gly Asn Leu Pro Leu Ser Ile Thr Leu Glu Gly Ser Asn Phe 20 25 30

Leu Ala Asn Gly His Pro Phe Leu Thr Glu Val Pro Glu Asn Ile Ile 35 40 45

Val Thr Pro Ser Pro Ile Asp Ala Lys Ser Ser Lys Asn Asn Glu Asp 50 55 60

Asp Asp Val Val Gly Cys Phe Val Gly Phe His Ala Asp Glu Pro Arg 65 70 75 80

Ser Arg His Val Ala Ser Leu Gly Lys Leu Arg Gly Ile Lys Phe Met $85 \hspace{1cm} 90 \hspace{1cm} 95$

Ser Ile Phe Arg Phe Lys Val Trp Trp Thr Thr His Trp Val Gly Ser 100 105 110

Asn Gly His Glu Leu Glu His Glu Thr Gln Met Met Leu Leu Asp Lys 115 120 125

Asn Asp Gln Leu Gly Arg Pro Phe Val Leu Ile Leu Pro Ile Leu Gln 130 135 140

Ala Ser Phe Arg Ala Ser Leu Gln Pro Gly Leu Asp Asp Tyr Val Asp 145 150 155 160

Val Cys Met Glu Ser Gly Ser Thr Arg Val Cys Gly Ser Ser Phe Gly 165 170 175

Ser Cys Leu Tyr Val His Val Gly His Asp Pro Tyr Gln Leu Leu Arg 180 185 190

Glu Ala Thr Lys Val Val Arg Met His Leu Gly Thr Phe Lys Leu Leu 195 200 205

Glu Glu Lys Thr Ala Pro Val Ile Ile Asp Lys Phe Gly Trp Cys Thr 210 220

Trp Asp Ala Phe Tyr Leu Lys Val His Pro Ser Gly Val Trp Glu Gly 225 230 235 240

Val Lys Gly Leu Val Glu Gly Gly Cys Pro Pro Gly Met Val Leu Ile 245 250 255

Asp Asp Gly Trp Gln Ala Ile Cys His Asp Glu Asp Pro Ile Thr Asp 260 265 270

Gln	Glu	Gly 275	Met	Lys	Arg	Thr	Ser 280	Ala	Gly	Glu	Gln	Met 285	Pro	Cys	Arg
Leu	Val 290	Lys	Leu	Glu	Glu	Asn 295	Tyr	Lys	Phe	Arg	Gln 300	Tyr	Cys	Ser	Gly
Lys 305	Asp	Ser	Glu	Lys	Gly 310	Met	Gly	Ala	Phe	Val 315	Arg	Asp	Leu	Lys	Glu 320
Gln	Phe	Arg	Ser	Val 325	Glu	Gln	Val	Tyr	Val 330	Trp	His	Ala	Leu	Cys 335	Gly
Tyr	Trp	Gly	Gly 340	Val	Arg	Pro	Lys	Val 345	Pro	Gly	Met	Pro	Gln 350	Ala	Lys
Val	Val	Thr 355	Pro	Lys	Leu	Ser	Asn 360	Gly	Leu	Lys	Leu	Thr 365	Met	Lys	Asp
Leu	Ala 370	Val	Asp	Lys	Ile	Val 375	Ser	Asn	Gly	Val	Gly 380	Leu	Val	Pro	Pro
His 385	Leu	Ala	His	Leu	Leu 390	Tyr	Glu	Gly	Leu	His 395	Ser	Arg	Leu	Glu	Ser 400
Ala	Gly	Ile	Asp	Gly 405	Val	Lys	Val	Asp	Val 410	Ile	His	Leu	Leu	Glu 415	Met
Leu	Ser	Glu	Glu 420	Tyr	Gly	Gly	Arg	Val 425	Glu	Leu	Ala	Lys	Ala 430	Tyr	Tyr
Lys	Ala	Leu 435	Thr	Ala	Ser	Val	Lys 440	Lys	His	Phe	Lys	Gly 445	Asn	Gly	Val
Ile	Ala 450	Ser	Met	Glu	His	Cys 455	Asn	Asp	Phe	Phe	Leu 460	Leu	Gly	Thr	Glu
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His	Cys	Ala	Tyr 500	Asn	Ser	Leu	Trp	Met 505	Gly	Asn ·	Phe	Ile	Gln 510	Pro	Asp
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Ser	Arg 530	Ala	Ile	Ser	Gly	Gly 535	Pro	Val	Tyr	Val	Ser 540	Asp	Cys	Val	Gly
Lys 545	His	Asn	Phe	Lys	Leu 550	Leu	Lys	Ser	Leu	Ala 555	Leu	Pro	Asp	Gly	Thr 560
Ile	Leu	Arg	Cys	Gln 565	His	Tyr	Ala	Leu	Pro 570	Thr	Arg	Asp	Cys	Leu 575	Phe
Glu	Asp	Pro	Leu 580	His	Asp	Gly	Lys	Thr 585	Met	Leu	Lys	Ile	Trp 590	Asn	Leu
Asn	Lys	Tyr	Thr	Gly	Val	Leu	Gly	Leu	Phe	Asn	Cys	Gln	Gly	Gly	Gly

595		600	605	
Trp Cys Pro Val	Thr Arg Arg 615		Ala Ser Glu P	he Ser Gln
Thr Val Thr Cys	Leu Ala Ser 630	Pro Gln Asp	Ile Glu Trp S	er Asn Gly 640
Lys Ser Pro Ile	Cys Ile Lys 645	Gly Met Asn 650	Val Phe Ala V	al Tyr Leu 655
Phe Lys Asp His 660	_	Leu Met Lys 665		ys Leu Glu 70
Val Ser Leu Glu 675	Pro Phe Thr	Phe Glu Leu 680	Leu Thr Val S 685	er Pro Val
Ile Val Leu Ser 690	Lys Lys Leu 695		Ala Pro Ile G 700	ly Leu Val
Asn Met Leu Asn 705	Thr Gly Gly 710	Ala Ile Gln	Ser Met Glu P 715	he Asp Asn 720
His Ile Asp Val	Val Lys Ile 725	Gly Val Arg 730	Gly Cys Gly G	lu Met Lys 735
Val Phe Ala Ser 740	Glu Lys Pro	Val Ser Cys 745	_	ly Val Val 50
Val Lys Phe Asp 755	Tyr Glu Asp	Lys Met Leu 760	Arg Val Gln V 765	al Pro Trp
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cag ttt cgg gct Gln Phe Arg Ala 35				
tgt gtc gaa agc Cys Val Glu Ser 50				3 3

										acc Thr				241
										ttc Phe				289
										Gly ggg				337
										gtt Val				385
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										gcg Ala				481
	-		_			-			-	ccc Pro	_	 _		529
										gag Glu				577
										ttt Phe				625
										gtg Val 220				673
										ccg Pro				721
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										ggc Gly				817
	_			_	_		_	_		tta Leu				865
										gtc Val 300				913

	atg Met															961
	tac Tyr	_	-			_		_								1009
	gtc Val															1057
	gag Glu															1105
	cca Pro 370															1153
	gtg Val															1201
	gat Asp		-	-			_				_	_	_			1249
	gcc Ala															1297
	gga Gly															1345
	tcc Ser 450			_	_	-					_		_	-	-	1393
	ttt Phe															1441
	tat Tyr															1489
	Gly															1537
	cac His															1585
	gga Gly 530															1633
tac	cta	ttc	cac	gag	aag	aaa	ctc	gtc	ctt	tct	aag	cca	tca	gac	aaa	1681

Tyr 545	Leu	Phe	His	Glu	Lys 550	Lys	Leu	Val	Leu	Ser 555	Lys	Pro	Ser	Asp	Lys 560	
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Lys	Ser	Asp	Asp 20	Arg	Pro	Tyr	Ile	Val 25	Leu	Leu	Pro	Leu	Ile 30	Glu	Gly	
Gln	Phe	Arg 35	Ala	Ser	Leu	Gln	Pro 40	Gly	Val	Asp	Asp	Phe 45	Ile	Asp	Ile	
Cys	Val 50	Glu	Ser	Gly	Ser	Thr 55	Lys	Val	Asn	Glu	Ser 60	Ser	Phe	Arg	Ala	
Ser 65	Leu	Tyr	Met	His	Ala 70	Gly	Asp	Asp	Pro	Phe 75	Thr	Leu	Val	Lys	Asp 80	
Ala	Val	Lys	Val	Ala 85	Arg	His	His	Leu	Gly 90	Thr	Phe	Arg	Leu	Leu 95	Glu	
Glu	Lys	Thr	Pro 100	Pro	Gly	Ile	Val	Asp 105	Lys	Phe	Gly	Trp	Cys 110	Thr	Trp	
Asp	Ala	Phe 115	Tyr	Leu	Asn	Val	Gln 120	Pro	His	Gly	Val	Met 125	Glu	Gly	Val	
Gln	Gly 130	Leu	Val	Asp	Gly	Gly 135	Cys	Pro	Pro	Gly	Leu 140	Val	Leu	Ile	Asp	
Asp 145	Gly	Trp	Gln	Ser	Ile 150	Cys	His	Asp	Asn	Asp 155	Ala	Leu	Thr	Thr	Glu 160	
Gly	Met	Gly	Arg	Thr 165	Ser	Ala	Gly	Glu	Gln 170	Met	Pro	Cys	Arg	Leu 175	Ile	
Lys	Phe	Glu	Glu 180	Asn	Tyr	Lys	Phe	Arg 185	Glu	Tyr	Glu	Ser	Pro 190	Asn	Lys	
Thr	Gly	Pro 195	Gly	Pro	Asn	Thr	Gly 200	Met	Gly	Ala	Phe	Ile 205	Arg	Asp	Met	
Lys	Asp 210	Asn	Phe	Lys	Ser	Val 215	Asp	Tyr	Val	Tyr	Val 220	Trp	His	Ala	Leu	

Cys Gly Tyr Trp Gly Gly Leu Arg Pro Asn Val Pro Gly Leu Pro Glu

225	230			235			240
Ala Lys Leu :	Ile Glu Pro 245	Lys Leu	Thr Pro 250	Gly Leu	Lys Thr	Thr 255	Met
Glu Asp Leu A	Ala Val Asp 260	Lys Ile	Val Asn 265	Asn Gly	Val Gly 270	Leu	Val
Pro Pro Glu I 275	Phe Val Glu	Gln Met 280	Tyr Glu	Gly Leu	His Ser 285	His	Leu
Glu Ser Val (290	Gly Ile Asp	Gly Val 295	Lys Val	Asp Val 300	Ile His	Leu	Leu
Glu Met Leu (305	Cys Glu Asp 310	Tyr Gly	Gly Arg	Val Asp 315	Leu Ala	Lys	Ala 320
Tyr Tyr Lys A	Ala Leu Ser 325	Ser Ser	Val Asn 330	Asn His	Phe Asn	Gly 335	Asn
Gly Val Ile A	Ala Gly Leu 340	Glu His	Cys Asn 345	Asp Phe	Met Phe 350	Leu	Gly
Thr Glu Ala 3	Ile Thr Leu	Gly Arg 360	Val Gly	Asp Asp	Phe Trp 365	Cys	Thr
Asp Pro Ser (Gly Asp Pro	Asn Gly 375	Thr Phe	Trp Leu 380	Gln Gly	Cys	His
Met Val His (385	Cys Ala Tyr 390	Asn Ser	Ile Trp	Met Gly 395	Asn Phe	Ile	His 400
Pro Asp Trp A	Asp Met Phe 405	Gln Ser	Thr His 410	Pro Cys	Ala Glu	Phe 415	His
Ala Ala Ser A	Arg Ala Ile 420	Ser Gly	Gly Pro 425	Ile Tyr	Val Ser 430	Asp	Ser
Val Gly Lys H 435	His Asn Phe	Glu Leu 440	Leu Arg	Ser Leu	Val Leu 445	Pro	Asp
Gly Ser Ile I 450	Leu Arg Cys	Asp Tyr 455	Tyr Ala	Leu Pro 460	Thr Arg	Asp	Cys
Leu Phe Glu <i>F</i> 465	Asp Pro Leu 470	His Asn	Gly Lys	Thr Met 475	Leu Lys	Ile	Trp 480
Asn Tyr Asn I	Lys Phe Thr 485	Gly Val	Val Gly 490	Thr Phe	Asn Cys	Gln 495	Gly
Gly Gly Trp S	Ser Arg Glu 500	Val Arg	Arg Asn 505	Gln Cys	Ala Ala 510	Glu	Tyr
Ser His Ala V 515	Val Ser Ser	Ser Ala 520	Gly Pro	Ser Asp	Ile Glu 525	Trp	Lys
Gln Gly Thr S 530	Ser Pro Ile	Asp Val 535	Asp Gly	Val Lys 540	Thr Phe	Ala	Leu
Tyr Leu Phe F 545	His Glu Lys 550	Lys Leu	Val Leu	Ser Lys 555	Pro Ser	Asp	Lys 560

Ile Asp Ile Thr Leu Glu Pro Phe Asp Phe Glu Leu Ile Thr Val Ser 565 570 575

Pro Val Lys Thr Leu Ala Asn Cys Thr Val 580 585

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	ggc Gly													97
	ctg Leu													145
	ggc Gly 50													193
	gac Asp													241
	gtc Val													289
	cgg Arg			_	_	-	-							337
	gcc Ala													385
	gtc Val 130													433
	agg Arg													481
_	cag Gln							_	_	_		_	_	529

	_				gcc Ala		_							-		577
					ggc Gly											625
					gca Ala											673
					gcg Ala 230											721
					tac Tyr											769
					tcg Ser											814
tgaç	accad	gac c	gggc	cgato	ga ct	ctg	cgtct	cto	gctco	cctg	ctg	gcct	gct d	cagga	acataa	874
tcta	atgt	tt a	agago	cttac	cc aç	ggttt	taca	a gct	ctat	cag	ttta	cttt	tg t	ctttt	ctgct	934
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Ser	Gly	Gly	Pro 20	Ile	Tyr	Val	Ser	Asp 25	Ser	Val	Gly	Gln	His 30	Asp	Phe	
Ala	Leu	Leu 35	Arg	Arg	Leu	Ala	Leu 40	Pro	Asp	Gly	Thr	Val 45	Leu	Arg	Cys	
Glu	Gly 50	His	Ala	Leu	Pro	Thr 55	Arg	Asp	Cys	Leu	Phe 60	Ala	Asp	Pro	Leu	

70

85

Gly Val Val Gly Ala Phe Asn Cys Gln Gly Gly Gly Trp Ser Pro Glu

Ala Arg Arg Asn Lys Cys Phe Ser Glu Phe Ser Val Pro Leu Ala Ala

Arg Ala Ser Pro Ser Asp Val Glu Trp Lys Ser Gly Lys Ala Gly Pro

Gly Val Ser Val Lys Asp Val Ser Gln Phe Ala Val Tyr Ala Val Glu 130 135 140	
Ala Arg Thr Leu Gln Leu Leu Arg Pro Asp Glu Gly Val Asp Leu Thr 145 150 155 160	
Leu Gln Pro Phe Thr Tyr Glu Leu Phe Val Val Ala Pro Val Arg Val 165 170 175	
Ile Ser His Glu Arg Ala Ile Lys Phe Ala Pro Ile Gly Leu Ala Asn 180 185 190	
Met Leu Asn Thr Ala Gly Ala Val Gln Ala Phe Glu Ala Lys Lys Asp 195 200 205	
Ala Ser Gly Val Thr Ala Glu Val Phe Val Lys Gly Ala Gly Glu Leu 210 215 220	
Val Ala Tyr Ser Ser Ala Thr Pro Arg Leu Cys Lys Val Asn Gly Asp 225 230 235 240	
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His Trp Val Gly
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Val Tyr Val Trp His Ala Leu Cys Gly Tyr Trp Gly Gly Val Arg Pro
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Leu Val Pro Pro
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Phe Ala Pro Ile Gly Leu Val Asn Met
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<223> n = inosine
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ttnaangtnt ggtggacnac ncantgggtn gg
                                                                    32
<210> 39
<211> 41
<212> DNA
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<223> n = inosine
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                                                                      44
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tgccancent entenatnat nacnaancen ggnggneane ence
                                                                      44
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	Description of Artificial Sequence:synthetic primer 5-F (from list 4)	
<400> tggatg	45 ggna anttnatnca nccngantgg ganatgtt	38
<210> <211> <212> <212> <213> <	38	
<222>	<pre>modified_base (1)(38) n = inosine</pre>	
	Description of Artificial Sequence:synthetic primer 5-RV (from list 4)	
<400>		38

<210> <211> <212> <213>	27		
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<210><211><211><212><213>	20		
<220> <223>	Description of Artificial primer 13.4 (from list 5)	Sequence:synthetic.	
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<210><211><211><212><213>	24		
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<210> <211> <212>	21		

<213>	Artificial Sequence	
<220>		
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yttrto	cytcr tanagraatt t	21
	•	
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	primer RES-2RV (from list 6)	
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ccaaat	ggta catattggct ccaaggttgt	30
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	primer RS-8 (from list 6)	
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<210> <211> <212> <213>	30	
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<210> <211> <212> <213>	30	
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<210> <211> <212> <213>	37	
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<210> 69 <211> 30 <212> DNA <213> Artificial Sequence	
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ggccacatnt tnacnarnce natnggngen aa
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NZZSZ II- INOSTIC	
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<210> <211> <212> <213>	30	
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<210> <211> <212> <213>	25	
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<210><211><211><212><213>	30	
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<210> <211> <212> <213>	33	
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primer RS-RV (from list 14)

<400> accttc	84 eccat acacettttg gatgaacett caa	33
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<210> <211> <212> <213>	36	
	Description of Artificial Sequence:XbaI-NotI-SacI linker (from Fig. 1)	
<400> tgataa	86 Leta gageggeege cacegeggtg gagete	36
<210><211><211><212><213>	34	
	Description of Artificial Sequence: XbaI-NotI-SacI linker (from Fig. 1)	
<400> tctaga	87 Ittat caaaataaaa actggaccaa agac	34